From: Sent: To: Subject:

Pak, Yong Tuesday, January 15, 2002 1:38 PM STIC-Biotech/ChemLib 09/654,652

dear stic,

please search the following in commercial and interference database for 09/654,652:

- 1. SEQ ID NO:3
- 2. oligomer search of SEQ ID NO:3

Yong Pak

Art Unit 1652 CM1 10A16 (mail box 10C01) 703-308-9363

Pending Nucleic Acid and/or Pending Amino Acid database searches now generate two sets of results. These databases were split into to two parts to reduce the time needed to update the databases daily. The split freed up more machine time for processing searches.

Searches run against the Nucleic Acid Pending database produce two sets of results, with the extensions, .rnpm and .rnpn

Searches run against the Amino Acid Pending database produce two sets of results, with the extensions, .rapm and .rapn

The Pending database search results should not be left in the case because they contain data that is confidential.

Searcher: Aleson Jaly
Phone: 308-450/
Location: Sinter lub.
Date Picked Up: 1/15/02
Date Completed: 1/15-102
Searcher Prep/Review:
Clerical: 5 2
Online time: 4

TYPE OF SEARCH:
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

VENDOR/COST(where applic.)
STN:
DIALOG:
Questel/Orbit:
DRLink:
Lexis/Nexis:
Sequence Sys.: ABSSO2
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Other (specify):

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Listing first 45 summaries
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Perfect score:
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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length: 2000000000
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US-09-286-690-10
US-09-286-690-7
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US-09-105-106-12
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RESULT 2
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; Sequence 2, Application US/08103998
; Patent No. 5470725

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GENERAL INFORMATION:
APPLICANT: Li, Xin-Liang
APPLICANT: Liungdahl, Lars G.
APPLICANT: Chen, Huizhong
TITLE OF INVENTION: Lichenase and Coding Sequences
FILE REFERENCE: 55-96
CURRENT APPLICATION NUMBER: US/09/286,690
CURRENT FILING DATE: 1999-04-05
                                                                                                                                                           Sequence 8, Application US/09286690 Patent No. 6103511
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Best Local Similarity
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SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: FOLEY & LARDNER
STREET: 1800 Diagonal Road, P.O. Box 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 30 TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 836-9300
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TITLE OF INVENTION: A Thermostable (1,3-1,4)-beta-Glucanase
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 16-OCT-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bent, Stephen A.
REGISTRATION NUMBER: 29,
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APPLICATION NUMBER: US 07/773,652
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TYPE: amino acid
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TELEX: 899149
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                                                                                                                                                                                                                                                                               KHT---ATANIPSTPGKIMMNLWNGTGVDDWLGSYNGAN-PLYAEYDWV---KYT
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Hofemeister, Ju
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Olsen, Ole
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SYSTEM: PC-DOS/MS-DOS
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; TYPE: PRT
; ORGANIZM: Bacillus subtilis
US-09-286-690-8
                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Bacillus licheniformis US-09-286-690-10
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US-09-286-690-10
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SEQ ID NO 10
LENGTH: 243
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Best Local Similarity
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Best Local Similarity
Matches 63; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: L1, Xin-Liang
APPLICANT: L1, Xin-Liang
APPLICANT: Chen, Huizhong
APPLICANT: Chen, Huizhong
TITLE OF INVENTION: Lichenase and Coding Sequences
FILE REFERENCE: 55-96
CURRENT APPLICATION NUMBER: US/09/286,690
CURRENT FILING DATE: 1999-04-05
EARLIER APPLICATION NUMBER: US 60/027,882
EARLIER APPLICATION NUMBER: US 60/027,882
EARLIER FILING DATE: 1999-10-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EARLIER APPLICATION NUMBER: PCT/US97/17811
EARLIER FILING DATE: 1997-10-03
NUMBER OF SEQ ID NOS: 12
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EARLIER FILING DATE: 1999-10-04
EARLIER APPLICATION NUMBER: PCT/US97/1781:
EARLIER FILING DATE: 1997-10-03
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 24
TYPE: PRT
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                                             143 RKTEGGQVSNLTGTQGLRFNLWSSESA-AWVGQFDESKLPLFQFINWVKVYK 193
  194
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                                                                                                                                              83 EVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQEV 142
                                                                                                                                                                                              81 TSPSYNKFDCGENRSVQTYGYGLYEVNMKPAKNVGIVSSFFTYTGPT---DGTPWDEIDI 137
                                                                                                                                                                                                                                            23 TNVSAKDFSGAELYTLEEVQYGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWVEVDI 82
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KHTATTQIPQTPGK--IMMNLWNGAGVDEWLGSYN-GVTPLSRSLHWVRYTK 242
                                                                                                EFLGKDTTKVQFNYYTNGVG---NHEKIVNLGFDAANSYHTYAFDWQPNSIKWYVDGQ-L 193
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                                                                                                                                                                                                                                                                                             12.8%; Score 233.5; DB 3; 33.7%; Pred. No. 1.1e-13; tive 26; Mismatches 77;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13.6%; Score 248; DB 3; Length 242; 36.0%; Pred. No. 5.3e-15; tive 25; Mismatches 75; Indels
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                                                                                                                                                                                                                                                                                                                                        DB 3;
                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                             243;
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                                                                                                                                                                                                                                                                                          Gaps
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RESULT

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TYPE: PRT
; ORGANISM: Clostridium thermocellum
US-09-286-690-9
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  В
                                                                                                                                                                            ; ORGANISM: Bacillus polymyxa
US-09-286-690-7
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                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 7
LENGTH: 238
TYPE: PRT
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                                                                              Matches
                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7, Application US/09286690 Patent No. 6103511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local Similarity
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Patent No. 610351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EARLIER APPLICATION NUMBER: US 60/027,882
EARLIER FILING DATE: 1999-10-04
EARLIER APPLICATION NUMBER: PCT/US97/17811
EARLIER FILING DATE: 1997-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/286,690 CURRENT FILING DATE: 1999-04-05
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                                                                                                                                                                                                                                                                                                                EARLIER APPLICATION NUMBER: US 60/027,882
EARLIER FILING DATE: 1999-10-04
EARLIER APPLICATION NUMBER: PCT/US97/17811
EARLIER FILING DATE: 1997-10-03
                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/286,690 CURRENT FILING DATE: 1999-04-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Li, Xin-Liang
APPLICANT: Ljungdahl, Lars G.
APPLICANT: Chen, Huizhong
TITLE OF INVENTION: Lichenase and Coding Sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 55-96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                196 VYR----GTRNIPVTPGKIMMNLWPGIGVDEWLGRYD-GRTPLQAEYEYVKYY 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     142 VRKTEGGQVSNLTGTQG-LRFNLWSSESA-AWVGQFDESKLPLFQFINWVKVY 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 82
  74
                                     20
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                  ALTTINVSAKDFSGAELYTLEEVQYGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWVE 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IEVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQE 141
SLTSPANNK-FDCGEYRSTNNYGYGLYEVSMKPAKNTGIVSSFFTYTGPSH---GTQWDE 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REYGGSYPYKSGEYRTKSFFGYGYYEVRMKAAKNVGIVSSFFTYTGPS---DNNPWDEID 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IEFLGKDTTKVQFNWYKNGVGG---NEYLHNLGFDASQDFHTYGFEWRPDYIDFYVDGKK 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61;
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                                                                              Conservative
                                                                                              12.5%;
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                                                                              26; Mismatches
                                                                            Score 227.5; DB 3;
Pred. No. 3.9e-13;
6; Mismatches 74;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 231.5; DB 3; Pred. No. 2.1e-13;
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                                                                              Indels
                                                                            17;
                                                                            Gaps
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; LENGTH: 242
; TENGTH: 242
; ORGANISM: Clostridium thermocellum
US-09-286-690-11
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EARLIER APPLICATION NUMBER: PCT/US97/17811
EARLIER FILING DATE: 1997-10-03
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 11
                                                                                                                                                                                                                                                             Sequence 4, Application US/08103998 Patent No. 5470725
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Li, Xin-Liang
APPLICANT: Ljungdahl, Lars G.
APPLICANT: Chen, Huizhong
                                                                                                                                                                                                                      GENERAL INFORMATION: APPLICANT: Borris:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/286,690 CURRENT FILING DATE: 1999-04-05 EARLIER APPLICATION NUMBER: US 60/027,882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Lichenase and Coding Sequences FILE REFERENCE: 55-96
                                                                                                                                                                           APPLICANT: APPLICANT:
                                                             CORRESPONDENCE ADDRESS:
                                                                                      TITLE OF INVENTION: A Thermostable NUMBER OF SEQUENCES: 4
                                                                                                                                                        APPLICANT:
                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                             178 SKLPL 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        172 DFHTYGFEWRPDYIDFYVDGKKVYR----GTRNIPVTPGKIMMNLWPGIGVDEWLGRYD-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             118 SSFFTYTGPS---DNNPWDEIDIEFLGKDTTKVQFNWYKNGVGG---NEYLHNLGFDASQ 171
                                                                                                                                                                                                                                                                                                                                                                                                 227 GRTPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               120 AFHTYGLEWTPNYVRWTVDGQEVRKTEGGQVSNLTGTQG-LRFNLWSSESA-AWVGQFDE 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14 VAAAAAALTTNVS------AKDFSGAELYTLEEVQ------YGKFEARMKMAAASGTV 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       80 VDIEVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDG 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60 SSMFLYQNGSEIADGRPWVEVDIEVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQ 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 58 VSTVLEAFTGDISNGKMILTLDREYGGSYPYKSGEYRTKSFFGYGYYEVRMKAAKNVGIV 117
                  ADDRESSEE: FOLEY & LARDNER STREET: 1800 Diagonal Road,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       QEVRKTEGGQVSNLTGTQG-LRFNLWSSESA-AWVGQFDESKLPLFQFINWVKVYKYT 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IDIEFLGKDTTKVQFNYYTNGVGGH----EKIINLGFDASTSFHTYAFDWQPGYIKWYVDG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
Alexandria
                                                                                                                                                                                                                                                                                                                                                                                                 231
                                                                                                                               Von Wettstein, Dietrich
                                                                                                                                                        Olsen, Ole
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                           Hofemeister, Jurgen
Thomsen, Karl Kristian
                                                                                                                                                                        Thomsen,
                                                                                                                                                                                                                      Borriss,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12.3%; Score 224.5; DB 3; 33.5%; Pred. No. 7.4e-13; mismatches 71;
                                                                                                                                                                                                                      Rainer
                      P.O.
                                                                                                           (1,3-1,4)-beta-Glucanase
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; ORGANISM: Orpinomyces sp. PC-2 US-09-286-690-2
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// MOLECULE TYPE: protein
US-08-103-998-4
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                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09286690 Patent No. 6103511
                                                                                SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/286,690 CURRENT FILING DATE: 1999-04-05
                                                                                                                                                                                                                                                                                                  APPLICANT: Li, Xin-Liang APPLICANT: Ljungdahl, Lars G. APPLICANT: Chen, Huizhong
                                                                                                                                                                              EARLIER APPLICATION NUMBER: US 60/027,882 EARLIER FILING DATE: 1999-10-04
                                                                                               SOFTWARE: PatentIn
                                                                                                                                      EARLIER APPLICATION NUMBER: PCT/US97/17811 EARLIER FILING DATE: 1997-10-03
                                                                                                                                                                                                                                                             TITLE OF INVENTION: Lichenase and Coding Sequences FILE REFERENCE: 55-96
                                                                                                                      NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 4:
                                         TYPE: PRT
                                                          LENGTH: 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 237 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 30 TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 836-9300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/773,652
FILING DATE: 16-CCT-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
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MEDIUM TYPE: FlOPPy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       188 KHTATTQIPAAPGK--IMMNLWNGTGVDDWLGSYNGVN-PIYAHYDWMRYRK 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          143 RKTEGGOVSNLTGTOGLRFNLWSSESA-AWVGOFDESKLPLFOFINWVKVYK 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       83 EVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQEV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23 TNVSAKDFSGAELYTLEEVQYGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWVEVDI 82
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EFLGKDTTKVQFNYYTNGAG----NHEKFADLGFDAANAYHTYAFDWQPNSIKWYVDGQ
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57; Conserv
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Pred. No. 1.1e-12;
Pred. No. 1.7matches 79;
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; TOPOLOGY: 11; MOLECULE TYPE: US-08-824-707-2
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                                                                                                                                    Query Match
Best Local Similarity
Matches 59; Conserv
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Best Local
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                                                                                                                                                                                                                                                                                                                          TELEFAX: 212-878-9655
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 42
TELECOMMUNICATION INFORMATION: 212-867-0123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 306 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION: NAME: Greqq, Valeta A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Savva, Demitris
TITLE OF INVENTION: No. 5919688el enzyme with beta-1,3-glucanase activity
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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163
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                                 60
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                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Gregg, Valeta A. REGISTRATION NUMBER: 35,127
                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 14
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A. ZIP: 10174-6401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           81 DIEVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQ 140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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SAFWMV--GANLPD-TPWPTSGEIDIMENVGNAPHEVHGTVHGPGYSGDNGIMGTYQHPQ 219
                               SSMFLYQNGSEIADGRPWV---EVDI-EVLGKNP----GSFQSNIITGKAGAQKTSEKHH 111
                                                                   NYTTSRVNSALDGQGNLVITALQESDGSYTSARLTTQGNVQPQFGRIEARIQIPRGQGIW
                                                                                                 NIKKTAVKSALAVAAAAALTTNVSAKDESGAELYTLEEV--QYGKFEARMKMAAASGTV 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AVYTA----YDNIPDTPGKIMMNAWNGIGVDDWLRPFN-GRTNISAYYDWV 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DIEFLGYDTTKVQFNYYTNGQGHH----EHIHYLGFDASQGFHTYGFFWARNSITWYVDGT 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LTIDRDGSGYTCGEYRTKNYYGYGMFQVNMKPIKNPGVVSSFFTYTGPS----DGTKWDEI 135
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                                                                                                                                      Conservative
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kington Avenue, Suite 6400
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27.6%;
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                                                                                                                                                     Score 159.5; DI
Pred. No. 9e-07;
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Pred. No. 9.6e-10;
                                                                                                                                      Mismatches
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                                                                                                                                                                     DB 2;
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                                                                                                                                                                    Length 306;
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LENGTH: 263
TYPE: PRT
                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                            Sequence 11, Application US/09159106 Patent No. 6284509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Applic
Patent No. 6284509
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                                                                                 APPLICANT: Ferrer, Pau
APPLICANT: Diers, Ivan
APPLICANT: Halkler, Torben
APPLICANT: Hedegaard, Lisbeth
                    TITLE OF INVENTION: An Enzyme With TITLE OF INVENTION: Activity FILE REFERENCE: 4693.204-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EARLIER APPLICATION NUMBER: 0885/96 EARLIER FILING DATE: 1996-08-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EARLIER APPLICATION NUMBER: 0427/96 EARLIER FILING DATE: 1996-12-04
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CURRENT FILING DATE: 1998-09-23
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CURRENT APPLICATION NUMBER: US/09/159,106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EARLIER FILING DATE: 1997-04-14
NUMBER OF SEQ ID NOS: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   42 QYGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWVEVDIEVLGKNPGSFQSNIITG-- 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 V--FDQ---PFFLILN-VAIGGQWPGNPDATTPF 289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ----KAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQEVRKTEGGQVSNLT 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QYGRIEARIQIPROQG-IWPAFWMLGGS--FPGTPWPSGEIDIM-ENVG-FEPHRVHGTV 136
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HGPGYSGGSGITGMYQHPQGWSFADTFHTFAVDWKPGEITWFVDGQQFHRVTRASVG--- 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VGQFDESKLPLFQFINWVKVYKYTPGQGEGGSDF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         GTQGLRFNLWSSESAAWVGQFDESKLPLFQFINWVKVYKYTPGQGEGGS----DFTLDWT 210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/09159106
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                                                                                                                                                                                                                                                                                                                               243
                                                                                                                                                                                                                                                                                                                                                                                                                 --ANAWV--FDQ---PFFLILN-VAVGGQWPGYPDGTTQLPQQMKVDYV 234
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 145; DB 4; 1
Pred. No. 1.5e-05;
9; Mismatches 73;
                                                               -1,3-Glucanase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 263;
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Query Match
Best Local Similarity
Thehes 78; Conserve
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US-09-159-106-11
US-09-159-106-13
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SOFTWARE: FRASTSEQ for Windows Version
SEQ ID NO 11
LENGTH: 435
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 6284509 GENERAL INFORMATION:
                                                                                        SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13,
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EARLIER APPLICATION NUMBER: 0427/96
EARLIER FILING DATE: 1996-12-04
EARLIER APPLICATION NUMBER: 0885/96
EARLIER FILING DATE: 1996-08-23
EARLIER APPLICATION NUMBER: PCT/DK97/00160
EARLIER FILING DATE: 1997-04-14
                                                                                                                                                           EARLIER APPLICATION NUMBER: 0885/96
EARLIER FILING DATE: 1996-08-23
EARLIER APPLICATION NUMBER: PCZ/DK97/00160
EARLIER FILING DATE: 1997-04-14
                                                                                                                                                                                                                                                   TITLE OF INVENTION: An Enzyme With -1,3-0
TITLE OF INVENTION: Activity
FILE REFERENCE: 4693.204-US
CURRENT APPLICATION NUMBER: US/09/159,106
CURRENT FILING DATE: 1998-09-23
EARLIER APPLICATION NUMBER: 0427/96
EARLIER FILING DATE: 1996-12-04
EARLIER FILING DATE: 1996-12-04
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Halkier, Torben APPLICANT: Hedegaard, Lisbeth
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                                                                                                              NUMBER OF SEQ ID NOS: SOFTWARE: FastSEQ for
                   ORGANISM: Oerskovia
                                            TYPE: PRT
                                                                  LENGTH: 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              148
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        314 RAANGMCVDVPWADPTDGNPVQIVTCSGNAAQTWTRGS---DGT-VRALGKCLDVRDG 367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   254 QPFFLILNVAVGGQWPGYPDGTTQLPQQMKVDYVRVYDNGSGSSSPGNPGTGLPTGTGAV 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 203 MYQHPQGWSFADTFHTFAVDWKPGEITWFVDGQQFHRVTRASVG------ANAWVFD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88 AELQNYTASRANSALDGQGNLVITARREGDGSYTSARMTTQGKYQPQYGRIEARIQIPRG 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        34 ---ELYTLE--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28 SALVAALTAAAAALAVTVAATSAAAAPGDLLWSDEFDGAAGSAPNPAVWNHETGAHGWGN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6 TAVKSALAVAAAAALT---TNVSA-----KDFSGA-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              QG-IWPAFWMLGGS--FPGTPWPSSGEIDIMENVGFEPHRVHGTVHGPGYSGGSGI--TG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SGTVSSMFLYQNGSEIADGRPWV---EVDI-EVLGKNP----GSFQSNIITGKAGAQKTS 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ------SAAWVGQFD-ESKLPLFQFINWVKVYKYTPGQGEGGSDFT------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Application US/09159106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -LDWTDNFD-----TFDGS---RWGKGDWTFDGNRVDLTDKNIYSRDG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.9%;
                      xanthineolytica
                                                                                                                 Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 144; DB 4;
Pred. No. 3.9e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                  TELEFAX: (212) 286-0854 or 286-0082
TELEX: TWX 710-581-4766
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH 1.44 Mb STORAGE DISKETTE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: MS-DOS
                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Bogosian, Elizabeth A.
REGISTRATION NUMBER: 39,911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: EN NUMBER OF SEQUENCES: 1 CORRESPONDENCE ADDRESS:
                 ORIGINAL SOURCE:
                                    FRAGMENT TYPE:
                                                 HYPOTHETICAL: I
                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 697-5995
                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 50 CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NERAL INFORMATION: APPLICANT: Jack G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  203 MYQHPQGWSFADTFHTFAVDWKPGEITWFVDGQQFHRVTRASVG-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   108 EKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQEVRKTEGGQVSNLTGTQGLRFNLWSSE
                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/OFILING DATE: 11-SEP-1996 CLASSIFICATION: 435
ORGANISM:
                                                                                                                                                                                                                                TELEPHONE:
                                                                                                     TOPOLOGY:
                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                             FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      56 SGTVSSMFLYQNGSEIADGRPWV---EVDI-EVLGKNP----GSFQSNIITGKAGAQKTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88 AELQNYTASRANSALDGQGNLVITARREGDGSYTSARMTTQGKYQPQYGRIEARIQIPRG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34 ---ELYTLE-----
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                                                                                                                                                   LENGTH:
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                                                                                                                                   amino acid
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gub, Rhodothermus marinus
                                                                                                  linear
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                                                                                  peptide
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23.0%; Pred. No. 3.
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RESULT 15
US-08-712-072C-3
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US-08-712-072C-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Jack Gold
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                           TELEX: TWX 710-581-4766 INFORMATION FOR SEQ ID NO:
          FRAGMENT TYPE:
ORIGINAL SOURCE:
ORGANISM: e13
                                                                            MOLECULE TYPE: pe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH 1.44 N
COMPUTER: IBM PC COMPATIBLE
                                                             HYPOTHETICAL: 1
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION: NAME: Bogosian, Elizabet
                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Allocation Avenue
                                                                                                                            TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 39,911
REFERENCE/DOCKET NUMBER: 63475/97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    83 -EVLGKNP----GSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTV 137
                                                                                                                                                                                                                           TELEFAX:
                                                                                                                                                                                                                                            TELEPHONE:
                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: ASCII
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93 REYTSARLVTRGKASWTYGRFEIRARLPSGRGTWPAIWMLPDRQTYGSAY-WPDNGEIDI 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28 KDFSGAELYTLEEVQ--YGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWV---EVDI 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
nes 44; Conserv
                                                                                                                                                            LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DGQEVRKTEGGQVSNLTG - -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10016
                                                                                                                                                                                                                                                                                                         Bogosian, Elizabeth A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ΝY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/08712072C
                                                                                                                                                         321 amino acids
                                                                                                                                                                                                                               (212)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jack Goldstein,
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             e13b,
                                                                                                              linear
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                                                                                                                                                                                                                             286-0854 or 286-0082
             Bacillus circulars
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23.7%;
                                                                                                                                                                                                                                                697-5995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INCH 1.44 Mb STORAGE DISKETTE
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ENDO-BETA-GALACTOSIDASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rothstein & Ebenstein
                                                                                                                                                                                                                                                                                                                                                                                                                             US/08/712,072C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ---TQGLRFNLWSSESAAWVGQ--FDESKLPLFQFINW 188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 133.5;
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Search completed: January 15, 2002, 15:53:08 Job time: 288 sec

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